

B1

X_8 is any amino acid residue;
 X_9 is an aliphatic residue;
 X_{10} is any amino acid residue;
 Z_3 is an optional 1 to 5 residue peptide or peptide analog;
 Z_4 is $-C(O)OR$ or $-C(O)NRR$;
each R is independently hydrogen, (C_1-C_6) alkyl, (C_2-C_6) alkenyl, (C_2-C_6) alkynyl or (C_6-C_{14}) aryl;
each $"-"$ between residues X_1 through X_{10} , Z_2 and X_1 and X_{10} and Z_3 independently represents an amide linkage, a substituted amide linkage or an isostere of an amide linkage; and each $"\sim"$ represents a bond.

Please replace claim 12 with the following:

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12. (once amended) The compound of claim 1 which is a 10-20 residue peptide or peptide analog according to formula (I):

(I) $Z_1 \sim Z_2 \sim X_1 \sim X_2 \sim X_3 \sim X_4 \sim X_5 \sim X_6 \sim X_7 \sim X_8 \sim X_9 \sim X_{10} \sim Z_3 \sim Z_4$

or a pharmaceutically-acceptable salt thereof, wherein:

Z_1 is $R-C(O)-NR-$ or $RRN-$;

Z_2 is an optional 1 to 5 residue peptide or peptide analog;

X_1 is any amino acid residue;

X_2 is any amino acid residue;

X_3 is a hydrophobic residue or a hydroxyl-substituted aliphatic residue;

X_4 is any amino acid residue;

X_5 is a hydrophobic residue or Gly;

X_6 is a hydrophobic or a hydrophilic residue;

X_7 is Gly, an amido-substituted polar residue or a hydrophobic residue;

X_8 is any amino acid residue;

X_9 is an aliphatic residue;

X_{10} is any amino acid residue;

Z_3 is an optional 1 to 5 residue peptide or peptide analog;

Z_4 is $-C(O)OR$ or $-C(O)NRR$;

each R is independently hydrogen, (C_1-C_6) alkyl, (C_2-C_6) alkenyl, (C_2-C_6) alkynyl or (C_6-C_{14}) aryl;

each $"-"$ between residues X_1 through X_{10} , Z_2 and X_1 and X_{10} and Z_3 independently represents an amide linkage, a substituted amide linkage or an isostere of an amide linkage; and each $"\sim"$ represents a bond.